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SEQUENCE LISTING

0> Noteborn, Mathieu H.M.Astrid, Danen-Van Oorschot AAM

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TECH CENTER 1600/2900

<120> Apoptin-Associating Proteins

<130> 2906-4995US

<140> 09/655,109

<141> 2000-09-05

<150> EP 99202858.9

<151> 1999-09-02

<150> EP 99203465.2

<151> 1999-10-21

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<170> PatentIn version 3.1

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<213> Artificial Sequence

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17

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5

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15

<210> 4

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- <211> 947
- <212> DNA
- <213> Homo sapiens
- <220>
- <221> misc_feature .
- <222> (5)..(5)
- <223> N may be any nucleotide
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- <223> AAP-1-a nucleic acid
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- <221> misc feature
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- <223> N may be any nucleotide
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 teggateaat teteagetgg tggeneaaca agtggeacaa eagtatgeea eeceaceace 180
 eectaaaaag gagaagaagg agaaagttga aaageaggae aaagagaaac etgagaaaga 240
 eaaggaaatt agteetagtg ttaccaagaa aaataccaac aagaaaacca aaccaaagte 300

tgacattetg aaagateete etagtgaage aaacagcata cagtetgeaa atgetacaac 360

aaagaccage gaaacaaate acaceteaag geeeeggetg aaaaaacgtgg acaggageac 420

tgeacagcag ttggeagtaa etgtgggeaa egteacegte attateacag actttaagga 480

aaagactege teeteatega cateeteate cacagtgace teeagtgeag ggteagaaca 540

geagaaccag ascagetegg ggteagagag cacagacaag ggeteeteee gtteeteeac 600

geeaaaggge gacatgteag eagteaatga tgaatettte tgaaattgea catggaattg 660

tgaaaactat gaateagggt atgaaattea aaaceteeac etgeeeatge tgettgeate 720

cetggagaat ettetgtgga categacete ttagtgatge tgeeaggata atttetgett 780

geeatgggea tetggeeace aaggaattte geaceetgae gattaetett gacaetttta 840

tgtatteeat tgttttatat gatttteeta acaateattt ataattggat gtgeteetga 900

atetaetttt tataaaaaaaa geettygtgg eetegagaga tetatga 947

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<212> DNA

<213> Homo sapiens

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<223> AAP-1-b nucleic acid

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<213> Homo sapiens

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<222> (1)..(352)

<223> X is unknown amino acid residue

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Ala Gly Tyr Ser Pro Ser Met Thr Met Gly Asp Lys Lys Ser Pro Thr 20 25 30

Arg Pro Lys Arg Gln Ala Lys Pro Ala Ala Asp Glu Gly Phe Trp Asp 35 40 45

Cys Ser Val Cys Thr Phe Arg Asn Ser Ala Glu Ala Phe Lys Cys Ser 50 55 60

Ile Cys Asp Val Arg Lys Gly Thr Ser Thr Arg Lys Pro Arg Ile Asn 65 70 75 80

Ser Gln Leu Val Ala Gln Gln Val Ala Gln Gln Tyr Ala Thr Pro Pro 85 90 95

Pro Pro Lys Lys Glu Lys Us Glu Lys Val Glu Lys Gln Pro Lys Glu Lys Pro Glu Lys Asp Lys Glu Ile Ser Pro Ser Val Thr Lys Lys Asn Thr Asn Lys Lys Thr Lys Pro Lys Ser Asp Ile Leu Lys Asp Pro Pro Ser Glu Ala Asn Ser Ile Gln Ser Ala Asn Ala Thr Thr Lys Thr Ser Glu Thr Asn His Thr Ser Arg Pro Arg Leu Lys Asn Val Asp Arg Ser Thr Ala Gln Gln Leu Ala Val Thr Val Gly Asn Val Thr Val Ile Ile Thr Asp Phe Lys Glu Lys Thr Arg Ser Ser Ser Thr Ser Ser Ser Thr Val Thr Ser Ser Ala Gly Ser Glu Gln Gln Asn Gln Ser Ser Ser Gly Ser Glu Ser Thr Asp Lys Gly Ser Ser Ala Ser Ser Thr Pro Lys Gly Asp Met Ser Ala Val Asn Asp Glu Ser Phe Xaa Asn Cys Thr Trp Asn

Cys Glu Asn Tyr Glu Ser Gly Tyr Glu Ile Gln Asn Leu His Leu Pro

260

265

270

Met Leu Leu Ala Ser Leu Glu Asn Leu Leu Trp Thr Ser Thr Ser Xaa 275 280 285

Xaa Cys Cys Gln Asp Asn Phe Cys Leu Pro Trp Ala Ser Gly His Gln 290 295 300

Gly Ile Ser His Pro Asp Asp Tyr Ser Xaa His Phe Tyr Val Phe His 305 310 315 320

Cys Phe Ile Xaa Phe Ser Xaa Gln Ser Phe Ile Ile Gly Cys Ala Pro 325 330 335

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<213> Artificial Sequence

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<223> pACT-AAP-1b forward primer

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- <210> 8
- <211> 4.0
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- <213> Artificial Sequence
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- <223> pACT-AAP-1b reverse primer
- <400> 8
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- 40

- <210> 9
- <211> 35
- <212> DNA
- <213> Artificial Sequence
- <220>
- <223> pACT-AAP-1b forward primer
- <400> 9
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- <210> 10
- <211> 46
- <212> DNA

<213> Artificial Sequence

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<223> pACT-AAP-1b reverse primer

<400> 10

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